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Original Paper

DCP-LA Activates Cytosolic PKC_E by Interacting with the Phosphatidylserine Binding/Associating Sites Arg50 and Ile89 in the C2-Like Domain

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Key Words

DCP-LA • PKCɛ • C2-like domain • Phosphatidylserine binding/associating sites • Direct interaction

Abstract

Background/Aims: The linoleic acid derivative DCP-LA selectively and directly activates PKCE. The present study aimed at understanding the mechanism of DCP-LA-induced PKCE activation. Methods: Point mutation in the C2-like domain on PKCE was carried out, and each kinase activity was monitored in PC-12 cells using a föerster resonance energy transfer (FRET) probe with cyan fluorescent protein (CFP) and yellow fluorescent protein (YFP) at the N- and C-terminal ends of PKCE, respectively, or in the cell-free systems using a reversed phase highperformance liquid chromatography (HPLC). Intracellular PKCE mobilization was monitored in PC-12 cells using mRuby-conjugated PKCE. DCP-LA binding to PKCE was assayed using a fluorescein conjugated to DCP-LA at the carboxyl-terminal end (Fluo-DCP). Uptake of DCP-LA into cells was measured in PC-12 ells. Results: In the FRET analysis, DCP-LA decreased the ratio of YFP signal intensity/CFP signal intensity in PC-12 cells and in the cell-free kinase assay, DCP-LA increased area of phosphorylated PKC substrate peptide, indicating DCP-LAinduced PKC_E activation. These effects were significantly suppressed by replacing Arg50 and Ile89 by Ala or Asn in the C2-like domain of PKCE. In the fluorescent cytochemistry, DCP-LA did not affect intracellular PKCE distribution. In the cell-free binding assay, Fluo-DCP, that had no effect on the potential for PKCE activation, bound to PKCE, and the binding was inhibited only by mutating Ile89. Extracellularly applied DCP-LA was taken up into cells in a concentrationdependent manner. Although no activation was obtained in the cell-free kinase assay, the broad PKC activator PMA activated PKCE in PC-12 cells in association with translocation towards the cell surface, which was inhibited by mutating I89A. Conclusion: Unlike PMA DCP-LA activates cytosolic PKCE by binding to the phosphatidylserine binding/associating sites Arg50 and Ile89, possibly at the carboxyl-terminal end and the cyclopropane rings, respectively.

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Introduction

Accumulating evidence has shown that the linoleic acid (LA) derivative 8-[2-(2-pentylcyclopropylmethyl)-cyclopropyl]octanoic acid (DCP-LA) could improve dementia including Alzheimer's disease [1-8]. The effect is mainly attributed to selective and direct activation of PKC ε involving delivery of α 7 ACh receptor to the presynaptic membrane surface and release of neurotransmitters [9-12]. In our earlier study, DCP-LA-induced PKC ε activation was inhibited by adding phosphatidylserine (PS) [13]. This raises the possibility that DCP-LA activates PKC ε by interacting with the PS binding site on PKC ε . The accurate binding sites, however, have not been identified.

The present study was conducted to answer this issue. We show here that unlike the broad PKC activator phorbol 12-myristate 13-acetate (PMA) DCP-LA binds to Arg50 and Ile89 in the C2-like domain of PKCɛ and activates PKCɛ localized in the cytosol, but not at the plasma membrane. We also show that LA activates PKCɛ by binding to the same sites as DCP-LA.

Materials and Methods

DNA subcloning and site-directed mutagenesis

The plasmid containing the normal rat PKCɛ was constructed. For the PKCɛ mutants the amino acid residues in the C2-like domain of PKCɛ were replaced by Ala or Asn.

FRET analysis

The vector for FRET probe with CFP and YFP at the N- and C-terminal ends of PKC ε , respectively, was constructed and transfected into PC-12 cells using a Lipofectamine LTX-PLUS (Invitrogen, Carlsbad, CA, USA). Forty-eight hours later FRET monitoring was carried out with Zeiss LSM510 META inverted microscope (Oberkochen, Germany) by the method as described previously [14]. The CFP and YFT fluorescent signals were detected at an absorbance of 474 and 506 nm, respectively, using an excitation light of 458 nm. The FRET ratio (YFP signal intensity/CFP signal intensity) was calculated using an ImageJ software (National Institutes of Health, USA).

Fluorescent cytochemistry

PKCε, conjugated with the red fluorescent protein eqFP611, mRuby, was expressed into PC-12 cells, and intracellular PKCε mobilization was monitored with a confocal scanning laser microscope (Axiovert/LSM510; Carl Zeiss, Oberkochen, Germany).

Cell-free PKCɛ assay

In the cell-free systems, PKCɛ activity was assayed by the method previously described [13, 15]. Briefly, a synthetic PKC substrate peptide (10 μ M) was reacted with wild-type and mutant PKCɛ in a Ca²⁺- and PS-free solution containing DCP-LA, lipids or PMA at 30°C for 5 min. After loading on a reversed phase HPLC (LC-10ATvp, Shimadzu Co., Kyoto, Japan), a substrate peptide peak and a new product peak were detected at an absorbance of 214 nm. Areas for non-phosphorylated and phosphorylated PKC substrate peptide were measured, and phosphorylated substrate peptide (pmol/min) was used as an index of PKC activity. PKCɛ activities induced by DCP-LA, lipids and PMA were calculated by subtracting the basal PKCɛ activities in the presence of DMSO (Δ pmol/min).

PKCe binding assay

Fluo-DCP was constructed by the method previously described [16]. Briefly, wild-type and mutant PKCɛ were reacted with Fluo-DCP in the presence and absence of non-conjugated DCP-LA, and then, separated by blue native-polyacrylamide gel electrophoresis. The fluorescent signals were visualized using a FluoroPhoreStar3000 (Anatech, Tokyo, Japan).

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Assay of intracellular distribution of DCP-LA

After treatment with DCP-LA for 5 min, lysates from PC-12 cells were separated into the cytosolic and plasma membrane components. DCP-LA in each component was labeled with 9-anthryldiazomethane (ADAM)(Funakoshi, Tokyo, Japan) and then, the sample solution was loaded onto the HPLC system. ADAM was detected at an excitation wavelength of 365 nm and an emission wavelength of 412 nm using a fluorescence detector.

Statistical analysis

Statistical analysis was carried out using analysis of variance (ANOVA) followed by a Bonferroni correction.

Results

DCP-LA activates PKCe without translocation towards the cell surface

We initially monitored PKC ϵ activity in PC-12 cells using a FRET probe, which has CFP and YFP at the N- and C-terminal ends of PKC ϵ , respectively (Fig. 1A). A conformational change of PKC ϵ from the inactive form to the active form leads to an enhancement of the CFP signal intensity from the YFP signal intensity (Fig. 1A), and therefore, a decrease and an increase in the ratio (YFP signal intensity/CFP signal intensity) correspond to activation and inactivation of PKC ϵ , respectively (Fig. 1B). PMA, DCP-LA and LA significantly decreased the FRET ratio as compared with that for DMSO (Fig. 1C). This implies that PMA, DCP-LA and LA activate PKC ϵ in PC-12 cells.

PKC ε contains the PS binding/associating sites in the C2-like domain (Fig. 1D) [17]. PMA-induced PKC ε activation was significantly inhibited by replacing Trp23 (W23A), Arg50 (R50A) and Ile89 by Ala (I89A) (Fig. 1E). Similarly, activation of mutant PKC ε W23A (mPKC ε W23A), -R50A and -I89A induced by DCP-LA was significantly attenuated as compared with that for wild-type PKC ε (wPKC ε) (Fig. 1E). These results suggest that PMA and DCP-LA activate PKC ε by interacting with the PS binding/associating sites Trp23, Arg50 and Ile89 in the C2-like domain.

In the fluorescent cytochemistry in PC-12 cells expressing mRuby-conjugated PKCɛ, PMA translocated wPKCɛ to the cell surface, but otherwise such effect was not found with DCP-LA and LA (Fig. 1F). This suggests that PKCɛ activation does not parallel with PKCɛ translocation towards the cell surface. PMA did not affect intracellular mPKCɛI89A distribution, while mPKCɛW23A, -R50A and -Y91A were translocated towards the cell surface in a fashion similar to that for wPKCɛ (Fig. 1F).

DCP-LA activates PKCE by interacting with Arg50 and Ile89 in the C2-like domain

In the cell-free PKCɛ assay, DCP-LA-induced activation of mPKCɛR50A, -I89A and -I89N (replacement of Ile89 by Asn) significantly reduced as compared with that for wPKCɛ, but no significant reduction was found with mPKCɛW23A and other mPKCɛ with replacement of Arg26 (R26A), Arg32 (R32A), His85 (H85A), Asp86 (D86A), Pro88 (P88A), Gly90 (G90A), Tyr91 (Y91A), Asp92 (D92A) and Asp93 (D93A) by Ala (Fig. 2A).

Likewise, LA activated wPKCɛ to an extent similar to that for DCP-LA, and the activation was significantly reduced by mutating Arg50 and Ile89 (Fig. 2B). Another unsaturated free fatty acid oleic acid (OA) and the saturated free fatty acid stearic acid (SA) activated wPKCɛ, but to a lesser extent than that for DCP-LA or LA, and the activation was also reduced by mutating Arg50 and Ile89 (Fig. 2C, D). Surprisingly, no activation of wPKCɛ was obtained with PMA (Fig. 2E). This, in the light of the fact that cell-free PKCɛ assay was carried out in a PS-free solution, suggests that PS is required for PMA-induced PKCɛ activation. Dioleoyl-PS (DO-PS) still activated wPKCɛ, and the activation was significantly reduced by mutating Ile89 and Gly90, but not Arg50 (Fig. 2F). This suggests that PS binds to Ile89 and Gly90, thereby partially activating PKCɛ.





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Fig. 1. FRET analysis. (A) A schema of the FRET probe. (B) The relation between the fluorescent intensities and absorbance wavelengths (left panel) and the FRET ratio (YFP signal intensity detected at an absorbance of 506 nm/CFP signal intensity detected at an absorbance of 474 nm). (C) The YFP and CFP signals were monitored in PC-12 cells expressing the FRET probe 15 min after treatment with DMSO, PMA (100 nM), DCP-LA (100 nM) or LA (100 nM). Data represent the mean (\pm SEM) FRET ratio (YFP signal intensity/CFP signal intensity)(n=10 independent experiments). *P* values, ANOVA followed by a Bonferroni correction. *NS*, not significant. (D) The PS binding/associating sites in the C2-like domain of PKCɛ. (E) Data represent the mean (\pm SEM) activities of wild-type (WT) and mutant PKCɛ (Δ FRET ratio)(n=10 independent experiments). *P* values as compared with wPKCɛ activity, ANOVA followed by a Bonferoni correction. (F) Fluorescent cytochemistry in PC-12 cells expressing mRuby 15 min after treatment with DMSO, PMA (100 nM), DCP-LA (100 nM) or LA (100 nM). Scale bars, 20 µm. Note that similar results were obtained from 10 independent experiments.

DCP-LA directly binds to free PKCE

To obtain evidence for DCP-LA binding to PKCε, we constructed Fluo-DCP (Fig. 3A). Fluo-DCP produced a single fluorescent signal band at 87 kDa in the electrophoresed gel, corresponding to the molecule of PKCε, and the signal was attenuated or abolished by co-treatment or pretreatment with non-conjugated DCP-LA (Fig. 3A). The signal intensity for Fluo-DCP binding to mPKCεI89A and -I89N was significantly lower than that to wPKCε, and no significant effect was obtained with the other mutants including mPKCεR50A (Fig. 3B). In the cell-free kinase assay, Fluo-DCP activated wPKCε to an extent similar to that for DCP-LA (Fig. 3C). These results indicate that DCP-LA directly binds to Ile89 in the PS binding site, to







OA



Fig. 2. Cell-free PKC ϵ assay. Synthetic PKC substrate peptide (10 µM) was reacted with wild-type (WT) and mutant PKC ϵ (0.1 µg/mL) as indicated in the absence and presence of DCP-LA (100 µM)(A), LA (100 µM)(B), OA (100 µM)(C), SA (100 µM)(D), PMA (1 µM)(E) or DO-PS (100 µM)(F) for 5 min. Data represent the mean (± SEM) PKC ϵ activity (Δ pmol/min)(n=4 independent experiments). *P* values as compared with WT PKC ϵ activity, ANOVA followed by a Bonferonni correction.



Fig. 3. DCP-LA binding assay. (A) Fluo-DCP (1 mM) was reacted with electrophoresed PKCɛ in the absence and presence of non-conjugated DCP-LA (Co-DCP)(1 mM) or after pretreatment with non-conjugated DCP-LA (Pre-DCP)(1 mM). CBB, Coomassie brilliant blue staining. Note that similar results were obtained with 4 independent experiments. (B) Fluo-DCP (1 mM) was reacted with electrophoresed wild-type (WT) and mutant PKCɛ as indicated. Data represent the mean (\pm SEM) signal intensity relative to that for wPKCɛ (n=4 independent experiments). *P* values as compared with the signal intensity for wPKCɛ, ANOVA followed by a Bonferonni correction. (C) PKCɛ activities induced by DMSO (None), DCP-LA (100 µM) and Fluo-DCP (100 µM). Data represent the mean (\pm SEM) PKCɛ activity (pmol/min)(n=4 independent experiments). *P* values, ANOVA followed by a Bonferonni correction. Fluo-DCP (1 mM) was reacted with electrophoresed PKC γ (D) or PKC ς (E) in the absence and presence of non-conjugated DCP-LA (Co-DCP)(1 mM) or after pretreatment with non-conjugated DCP-LA (Pre-DCP)(1 mM).



А

7

DCP-LA







Fig. 4. Uptake of DCP-LA into cells. PC-12 cells were treated with DCP-LA at concentrations as indicated, and DCP-LA in the cytosolic and plasma membrane components was quantified with the HPLC systems. Data represent the mean (\pm SEM) DCP-LA concentration (pmol/µg protein)(n=4 independent experiments).

Fig. 5. Putative DCP-LA binding site in the C2-like domain of PKCε.



activate PKC ϵ . Fluo-DCP, on the other hand, produced no signal band binding to the classical PKC isozyme PKC γ and the atypical PKC isozyme PKC γ (Fig. 3 D, E). Collectively, these results interpret that DCP-LA directly and preferentially binds to PKC ϵ .

DCP-LA is taken up into cells

We finally examined whether extracellularly applied DCP-LA is taken up into cells. When extracellularly applied to PC-12 cells, cytosolic DCP-LA increased in a concentration (10 nM-1 μ M)-dependent manner in parallel with increased concentrations in the plasma membrane (Fig. 4). This suggests that DCP-LA is capable of activating PKC ϵ localized in the cytosol.

Discussion

PMA is shown to activate PKCε by binding to the C1 domain [18]. In the present study, PMA activated wPKCε in PC-12 cells, and the activation was inhibited by replacing Trp23,



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Arg50 and Ile89 by Ala, yet no activation of wPKCɛ was obtained with PMA in the cell-free kinase assay using a PS-free solution. PMA translocated wPKCɛ towards the cell surface in PC-12 cells and the effect was cancelled in mPKCɛ189A. These results, in the light of the finding that PS partially activates PKC by interacting with Ile89 and Glv90, raise the possibility that an inactive form of PKC binds to PS in the plasma membrane at Ile89, to make PKC an open frame, allowing PMA to bind to the C1 domain and to activate PKC fully; in other words, PS is indispensable for PMA-induced PKC_ε activation. Extracellularly applied PMA, therefore, would be trapped in the plasma membrane, where PMA binds to the C1 domain of PKC_ɛ linked by PS (Fig. 6). Likewise, diacylglycerol, which is produced by phospholipase C-catalyzed hydrolysis of phosphatidylinositol 4,5-bisphosphate in vivo, may also activate PKC_ɛ at the plasma membrane.

Unlike PMA DCP-LA activated PKCE in PC-12 cells without translocation towards the cell surface. This suggests that DCP-LA activates PKCe localized in the cytosol. In support of this notion, DCP-LA was actually taken up into cells. DCP-LA-induced activation of PKCE in PC-12 cells and under the cell-free conditions was suppressed by replacing Arg50 and Ile89 by Ala or Asn. This implies that DCP-LA activates PKC by interacting with Arg50 and Ile89. Intriguingly, Fluo-DCP binding to PKCE was attenuated only by mutating Ile89, but not Arg50, although there was no significant difference in the potential of wPKC_ε activation between Fluo-DCP and DCP-LA. This, in the light of the fact that fluorescein is conjugated to DCP-LA at the carboxyl-terminal end, suggests that the carboxyl-terminal end on DCP-LA recognizes and binds to Arg50 on PKC_E. Overall, the results presented here lead to a conclusion that extracellularly applied DCP-LA is taken up into cells, where DCP-LA activates PKC_ɛ by binding to Arg50 and Ile89 in the C2-like domain, possibly at the carboxyl-terminal end and the cyclopropane rings, respectively (Fig. 5, 6).

Like DCP-LA LA activated PKCE without translocation, and the activation was also inhibited by mutating Arg50 and Ile89. This suggests that LA activates PKC by binding to Arg50 and Ile89 in the C2-like domain, possibly at the carboxyl-terminal end and the *cis*double bonds, respectively. The results of the present study, thus, may represent novel insight into the mechanism of PKCe activation induced by *cis*-unsaturated free fatty acids as well as DCP-LA.

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Disclosure Statement

The authors declare no competing financial interests.

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